

# Raw Sequence Listing Error Summary

FILE

#9  
Bill

## ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/383,894A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE 4-10-01

- 1 ☐ Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 ☐ Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 ☐ Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 ☐ Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 ☐ Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.  
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 ☐ Variable Length Sequence(s) \_\_\_\_\_ contain n's or Xaa's which represented more than one residue.  
As per the rules, each n or Xaa can only represent a single residue.  
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 ☐ PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) \_\_\_\_\_. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 8 ☐ Skipped Sequences (OLD RULES) Sequence(s) \_\_\_\_\_ missing. If intentional, please use the following format for each skipped sequence:  
(2) INFORMATION FOR SEQ ID NO:X:  
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")  
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:  
This sequence is intentionally skipped  
  
Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 ☐ Skipped Sequences (NEW RULES) Sequence(s) \_\_\_\_\_ missing. If intentional, please use the following format for each skipped sequence.  
<210> sequence id number  
<400> sequence id number  
000
- 10 ☒ Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 ☐ Use of <213>Organism (NEW RULES) Sequence(s) \_\_\_\_\_ are missing this mandatory field or its response.
- 12 ☐ Use of <220>Feature (NEW RULES) Sequence(s) \_\_\_\_\_ are missing the <220>Feature and associated headings.  
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial Sequence" or "Unknown"  
Please explain source of genetic material in <220> to <223> section.  
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 ☐ PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).  
Instead, please use "File Manager" or any other means to copy file to floppy disk.

1635

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/383,894A

DATE: 03/30/2001  
TIME: 15:37:27

Input Set : A:\00400191.app  
Output Set: N:\CRF3\03302001\I383894A.raw

3 <110> APPLICANT: Li, Ming  
5 <120> TITLE OF INVENTION: T-Type Calcium Channel  
7 <130> FILE REFERENCE: 004.00191  
9 <140> CURRENT APPLICATION NUMBER: US 09/383,894A  
10 <141> CURRENT FILING DATE: 1999-08-26  
12 <150> PRIOR APPLICATION NUMBER: US 60/098,004  
13 <151> PRIOR FILING DATE: 1998-08-26  
15 <150> PRIOR APPLICATION NUMBER: US 60/117,399  
16 <151> PRIOR FILING DATE: 1999-01-27  
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Does Not Comply  
Corrected Diskette Needed

## RAW SEQUENCE LISTING

DATE: 03/30/2001

PATENT APPLICATION: US/09/383,894A

TIME: 15:37:27

Input Set : A:\00400191.app

Output Set: N:\CRF3\03302001\I383894A.raw

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154 &lt;400&gt; SEQUENCE: 2

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156 1 5 10 15

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159 20 25 30

161 Gly Pro Gly Ser Thr Glu Lys Asp Pro Gly Ser Ala Asp Ser Glu Ala

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Input Set : A:\00400191.app

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171          85          90          95
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174          100          105          110
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177          115          120          125
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180          130          135          140
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183 145          150          155          160
185 Ile Ala Gly Met Leu Glu Tyr Ser Leu Asp Leu Gln Asn Val Ser Phe
186          165          170          175
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189          180          185          190
191 Arg Val Pro Ser Met Arg Ile Leu Val Thr Leu Leu Leu Asp Thr Leu
192          195          200          205
194 Pro Met Leu Gly Asn Val Leu Leu Leu Cys Phe Phe Val Phe Phe Ile
195          210          215          220
197 Phe Gly Ile Val Gly Val Gln Leu Trp Ala Gly Leu Leu Arg Asn Arg
198 225          230          235          240
200 Cys Phe Leu Pro Glu Asn Phe Ser Leu Pro Leu Ser Val Asp Leu Glu
201          245          250          255
203 Pro Tyr Tyr Gln Thr Glu Asn Glu Asp Glu Ser Pro Phe Ile Cys Ser
204          260          265          270
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207          275          280          285
209 Arg Gly Glu Gly Gly Gly Gly Pro Pro Cys Ser Leu Asp Tyr Glu Thr
210          290          295          300
212 Tyr Asn Ser Ser Ser Asn Thr Thr Cys Val Asn Trp Asn Gln Tyr Tyr
213 305          310          315          320
215 Thr Asn Cys Ser Ala Gly Glu His Asn Pro Phe Lys Gly Ala Ile Asn
216          325          330          335
218 Phe Asp Asn Ile Gly Tyr Ala Trp Ile Ala Ile Phe Gln Val Ile Thr
219          340          345          350
221 Leu Glu Gly Trp Val Asp Ile Met Tyr Phe Val Met Asp Ala His Ser
222          355          360          365
224 Phe Tyr Asn Phe Ile Tyr Phe Ile Leu Leu Ile Ile Val Gly Ser Phe
225          370          375          380
227 Phe Met Ile Asn Leu Cys Leu Val Val Ile Ala Thr Gln Phe Ser Glu
228 385          390          395          400
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231          405          410          415
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Output Set: N:\CRF3\03302001\I383894A.raw

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273 625           630           635           640
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278 Ala Asp Ser Gly Ala Cys Gly Pro Asp Ser Cys Pro Tyr Cys Ala Arg
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281 Thr Gly Ala Gly Glu Pro Glu Ser Ala Asp His Val Met Pro Asp Ser
282           675           680           685
284 Asp Ser Glu Ala Val Tyr Glu Phe Thr Gln Asp Ala Gln His Ser Asp
285       690           695           700
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Sequence

<400> 6  
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→ see item 10 on  
Envi Summary  
sheet

**VERIFICATION SUMMARY**

DATE: 03/30/2001

PATENT APPLICATION: US/09/383,894A

TIME: 15:37:28

Input Set : A:\00400191.app

Output Set: N:\CRF3\03302001\I383894A.raw

L:1216 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:6  
L:1216 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6  
L:1216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6